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(54) Title: NOVEL PROTEINS AND CLONED GENES FOR DIAGNOSIS AND PROPHYLAXIS OF BABESIOSIS

(57) Abstract

The subject invention concerns the identification of novel merozoite surface proteins of Babesia bovis. Also disclosed are monoclonal antibodies to these proteins as well genes which encode for the proteins. The invention further concerns the use of the novel proteins, recombinant DNA clones, and monoclonal antibodies in the detection, treatment, and prophylaxis of babesiosis.

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DESCRIPTION

NOVEL PROTEINS AND CLONED GENES FOR DIAGNOSIS AND PROPHYLAXIS OF BABESIOSIS

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Cross-Reference to a Related Application

This application is a continuation-in-part of our co-pending application Serial No. 333,155, filed April 4, 1989.

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Background of the Invention

Bovine babesiosis is a tick-transmitted, hemoparasitic disease caused by intraerythrocytic protozoa belonging to the genus <u>Babesia</u>. The disease caused by <u>Babesia</u> manifests itself clinically by fever and extensive hemolytic anemia that often leads to hypotensive shock, cerebral involvement, and death. More than a half billion cattle are estimated to be at risk of acquiring babesiosis. This disease represents a primary impediment to food and fiber production in much of the world.

To date, control of bovine babesiosis in enzootic areas has been partially successful through vaccination with attenuated strains of <u>Babesia</u> spp. or with more virulent strains followed by chemotherapeutic control. Protective immunity in babesiosis may be directed against one or more surface antigens associated with sporozoites, infected erythrocytes, and/or merozoites. Merozoite surface antigens are important in the pathogenesis and immunology of babesiosis due to their role in the parasite's recognition of, attachment to, and penetration of host erythrocytes and their accessibility to the immune system.

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Recently, progress has been made toward the identification and characterization of specific immunogens of merozoites of <u>Babesia bovis</u> (Smith, R.D., M.A. James, M. Ristic, M. Aikawa, and C.A. Vega Y Murgula [1981] Science 212:335-338; Wright, I.G., B.V. Goodger, K. Rode-Bramanis, J.S. Matlick, D.F. Mahoney, and D.J. Waltisbuhl [1983] Z. Parasitenkd. 69:703-714; Wright, I.G., G.B.

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Mirre, K. Rode-Bramanis, M. Chamberlain, B.V. Goodger, and D.J. Mahoney [1985] Infect. Immun. 48:109-113; Commins, M.A., B.V. Goodger, and I.G. Wright [1985] Int. J. Parasitol. 15:491-495; Wright, I.G. and P.W. Riddles [1986] "Biotechnological Control of Tick-Borne Disease," Meeting of the Food and Agriculture Organization of the United Nations, 6-10 October 1986, pp. 1-21, Rome, Italy; Waltisbuhl, D.J., B.V. Goodger, I.G. Wright, G.B. Mirre, and M.A. Commins [1987] Parasitol Res. 73:319-323; Goff, W.L., W.C. Davis, G.H. Palmer, and T.C. McGuire [1988] Infect. Immun. 56:2363-2368) and Babesia bigemina (McElwain, T.F., L.F. Perryman, W.C. Davis, and T.C. McGuire [1987] J. Immunol. 138(7):2298-2304). However, in only one instance (Smith et al., 1981) were antigens which provided protection against infection determined to be surface-exposed on merozoites as opposed to cytoplasmic in location.

Bovine babesiosis can be caused by either <u>Babesia bigemina</u> or <u>Babesia bovis</u>. These parasites have antigenic similarities and differences that may have important functional roles in the induction of protective immunity and antibody-based diagnosis. Also, <u>B. bovis</u> isolates, including the current Australian vaccine strain, are now known to consist of subpopulations that vary antigenically, in virulence, and in abundance within an isolate (Cowman, A.F., P. Timms, and D.J. Kemp [1984] Mol. Biochem. Parasitol. 11:91-103; Gill, A.C., A.F. Cowman, N.P. Stewart, D.J. Kemp, and P. Timms [1987] Exp. Parasitol. 63:180-188).

Current vaccine strategies include the use of attenuated live <u>Babesia bovis</u> parasites and various inactivated preparations (Montenegro-James, S., M. Toro Benitez, E. Leon, R. Lopez, and M. Ristic [1987] Parasitol. Res. 74:142-150; Smith et al., 1981; U.S. Patent No. 4,762,711 issued to Buening et al.; Kuttler, K.L., M.G. Levy, M.A. James, M. Ristic [1982] Am. J. Vet. Res. 43(2):281-284). The attenuated vaccine provides the best protection against challenge with both homologous and heterologous strains, although there are a number of serious disadvantages, including a short shelf-life, variation in virulence, contamination with host erythrocyte stroma, and perpetuation of the life cycle by creation of a carrier state. Inactivated vaccines induce protection against challenge with homologous

strains; however, only partial protection occurs against challenge with heterologous strains.

Animals that survive natural field infection or that recover from infection with an attenuated vaccine strain are protected against clinical disease. However, premunization in this manner is expensive, impractical in developing countries that lack the necessary infrastructure, and a potential mode of transmission for other blood-borne diseases.

Brief Summary of the Invention

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Disclosed and claimed here are novel merozoite proteins of <u>Babesia bovis</u>. These proteins are known to be expressed on the surface of the merozoite and may be used to raise neutralizing antibodies. Thus, they can be used in the formulation of subunit vaccines for the prophylaxis of bovine babesiosis. Several of the proteins described here raise antibodies to both <u>Babesia bovis</u> and <u>Babesia bigemina</u>, while others are species, or even isolate, specific.

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Also disclosed are monoclonal antibodies to bovine babesiosis antigens. These monoclonal antibodies are used to identify merozoite surface antigens and may be used in the treatment and/or diagnosis of bovine babesiosis.

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A further element of the invention is the identification of genes which code for <u>Babesia</u> proteins. These genes can be used to make recombinant proteins which can be utilized for vaccines.

The invention also provides a means of detecting the presence of disease-causing <u>Babesia</u> organisms. The detection method involves the use of DNA probes which selectively identify the presence of these organisms.

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Brief Description of the Drawings

Figure 1 is the translated DNA and amino acid sequence of lambda-Bo44.

Figure 2 is the DNA and amino acid sequence for rBv42.

Figure 3 is the DNA sequence for rBv60.

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Figure 4 is the amino acid sequence for rBv60.

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Detailed Description of the Invention

The subject invention pertains to the identification of surface-exposed proteins of <u>B. bovis</u> merozoites. The proteins of the invention have sizes of 16, 25, 37, 42, 44, 55, 60, 85, 98, 125, 145, 225, and 250 kDa. The evidence that the proteins are surface exposed includes: (i) monoclonal antibody binding of live merozoites, (ii) labeling by surface iodination, and (iii) sensitivity to mild trypsinization.

We have identified <u>B. bovis</u> merozoite proteins that, by virtue of their surface location and their reactivity with immune bovine sera, are candidates for subunit vaccines. Among the numerous proteins recognized by immune bovine sera, six proteins (37, 42, 55, 85, 125, and 145 kDa) appeared to be relatively immunodominant.

The 145 kDa protein was of parasite origin, but its location on the membrane surface was not directly apparent. This protein may have a small portion exposed at the surface of the merozoite that is sensitive to mild trypsinization but the epitope recognized by the monoclonal antibody located internally.

Immunoprecipitation of radiolabeled antigens with bovine antisera indicated that many <u>Babesia bovis</u> merozoite proteins contain isolate-common epitopes, while at least 8 <u>B. bovis</u> proteins contain species-cross-reactive epitopes. The amino acid sequence of three of the immunogenic proteins from <u>B. bovis</u> (42 kDa) have been determined. Amino acid sequences which deviate in insignificant ways from the disclosed amino acid sequences fall within the scope of the subject invention so long as the antigenic properties of the protein are not altered. Thus, the subject invention includes mutants and fragments of the amino acid sequences depicted herein which do not alter the protein secondary structure, or if the structure is altered, the antigenic activity is retained. In particular, it should be understood that conservative substitutions of amino acids may be made. For example, amino acids may be placed in the following classes: basic, hydrophobic, acidic, polar, and amide. Substitutions whereby an amino acid of one class is replaced with another amino

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acid of the same type fall within the scope of the subject invention so long as the substitution does not materially alter the antigenic activity of the compound.

The ability of antibodies against heterologous geographic isolates to immunoprecipitate proteins from the Mexico B. bovis isolate indicates the conservation of at least one and probably more epitopes between proteins from the heterologous isolates. The conservation of these epitopes is extensive, as many Mexico B. bovis isolate proteins were precipitated by antisera against a different geographic isolate (Honduras). The 42,000 molecular weight Mexico B. bovis protein was precipitated by all five of the undiluted and three of the diluted Honduras antisera.

Among the highly immunogenic <u>B. bovis</u> proteins, only one (the 42,000 molecular weight protein) is both isolate common and species specific. This protein can be used as an antigen for species-specific, antibody-based diagnosis.

Monoclonal antibodies (MoAbs) were generated against surface-exposed proteins on merozoites of <u>B. bovis</u>. A genomic library constructed in the lambdagt11 expression vector was screened with MoAbs for identification of clones expressing recombinant surface proteins. Four recombinant clones were identified.

Southern blot analyses confirmed the parasite-specificity of the cloned inserts. Western blot analyses demonstrated that recombinant protein production in these clones is IPTG-induced and that the recombinant molecules exist as beta-galactosidase fusion proteins.

Additionally, recombinant proteins, partially purified by affinity column chromatography and gel filtration chromatography, reacted with specific MoAbs in Western blot assay indicating that the integrity of the epitopes is retained during purification. Calves immunized with these partially purified recombinant proteins developed titers of between 10⁻² and 10⁻⁵ as evidenced by IFA-live. Immune sera from these animals immunoprecipitated metabolically-radiolabeled merozoite proteins confirming that determinants found on native proteins are expressed by the clones.

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DNA probe candidates were also identified using the lambda-gt11 genomic library of <u>B. bovis</u>. Two DNA sequences, designated lambda-Bo6 and lambda-Bo25, hybridized to <u>Babesia</u> DNA but not to bovine DNA. Bo6 detected Mexico and Australia isolates of <u>B. bovis</u> as well as <u>B. bigemina</u> DNA. Lambda-Bo25 demonstrated greater specificity; it did not hybridize detectably to <u>B. bigemina</u> DNA and showed greater sensitivity for Mexico isolates of <u>B. bovis</u> than for Australia isolates.

Thus, lambda-Bo6 is a good candidate for detecting <u>Babesia</u> infections in cattle and ticks, and lambda-Bo25 can be used to distinguish geographic isolates of <u>B. bovis</u>.

Materials and Methods

Strain of B. bovis, Stabilate Preparation, Cryopreservation, and In Vitro <u>Cultures</u>. The strain of <u>B. bovis</u> used in experiments outlined herein was originally isolated from a Boophilus microplus tick-induced infection in Mexico by Dr. R.D. Smith, University of Illinois at Urbana (Goff, W.L. and C.E. Yunker [1986] Exp. Parasitol. 62:202-210). The cloned line was derived from the Mexico isolate by limiting dilution cloning as previously described by Rodriguez et al. (Rodriguez, S.D., G.M. Buening, T.J. Green, and C.A. Carson [1986] Infect. Immun. 42:15-18). The parasites have been maintained in our laboratory by either repeated passages in splenectomized Holstein-Freisian bull calves or in vitro cultivation. Cryopreservation of stabilates of B. bovis-infected erythrocytes obtained from infected calves and the preparation of partially purified merozoites from thawed stabilates has been described (Palmer, D.A., G.M. Buening, and C.A. Carson [1982] Parasitol 84:567; McElwain, T.F., L.E. Perryman, W.C. Davis, and T.C. McGuire [1987] J. Immunol. 138:2298-2304). Continuous in vitro cultivation of B. bovis was performed using a modification of the microaerophilous stationary phase (MASP) culturing system (Goff and Yunker, 1986). Viability of merozoites obtained from either frozen stabilates or in vitro cultivation was confirmed by 6-CFDA assay (McElwain et al., 1987) prior to their use in experiments or immunizations.

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Isolation of Merozoites. Merozoites were harvested from cultures after the relative percentage of parasitized erythrocytes was increased by sequential reduction of the concentration of erythrocytes (Goff and Yunker, 1986). For collection of merozoites, the contents of flasks containing >15% parasitized erythrocytes were centrifuged at 400 x g for 10 min at 4°C. The supernatant was centrifuged at 3,000 x g for 15 min at 4°C to pellet the merozoites. The merozoites were suspended in Puck saline-glucose (saline-G), and 2 ml was overlaid on 10 ml of a preformed continuous gradient of 65% Percoll-35% Puck saline-G. The gradient was centrifuged in a swinging bucket rotor at 3,000 x g for 20 min at 4°C. The merozoites were isolated from a band with an approximate density of 1.069 g/ml between erythrocyte ghosts at the Percoll-Puck saline-G interface and the residual intact erythrocyte pellet. The merozoites were washed once in 0.15 M NaCl containing 0.01 M sodium citrate (CS), suspended in CS, and stored on ice until used (within 2 to 4 hr).

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Purification, Quantitation and Viability Estimation of Merozoites. An equal volume of the isolated merozoite suspension was mixed with 6-carboxy fluorescein diacetate (6-CFDA; final concentration in CS, 10 ug/ml; Calbiochem-Behring, La Jolla, CA) (McElwain et al., 1987). The mixture was incubated at room temperature for 20 min, followed by centrifugation at 1,000 x g for 10 min and was then suspended in phosphate-buffered saline (PBS; 0.15 M, pH 7.2) for counting on a hemacytometer. The sample was examined with phase microscopy and epifluorescence with a 40X oil objective and fluorescein filter (450 to 520 nm). Viability was assessed as the percentage of total merozoites emitting fluorescence.

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Preparation of Immune Bovine Sera. Two spleen-intact Holstein-Freisian steers, 14 months of age and indirect fluorescent-antibody test negative for <u>B. bovis</u>, <u>B. bigemina</u>, and <u>Anaplasma marginale</u>, were inoculated intravenously with approximately 6 x 10⁸ <u>B. bovis</u>-infected erythrocytes from the same blood stabilate used to initiate in vitro cultures. On day 9 postinoculation each steer developed detectable parasitemia and a febrile response which persisted through day 13 postinoculation. Antibody specific for <u>B. bovis</u> was detected with the indirect

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fluorescent-antibody test on day 10 postinoculation. The steers were challenge inoculated as before on days 48 and 80 postinoculation, and although the animals did not develop a fever or parasitemia, the antibody titer increased after each challenge. Sera were collected and stored at -70°C after the final challenge, when the indirect fluorescent-antibody test titer was 1:10,000.

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In addition, the cloned line was passed through a splenectomized calf whose blood at peak parasitemia was used to infect five 4-5 month old Holstein steers (5 x 10⁷ infected erythrocytes each) and to initiate in vitro cultures. The cattle were reinfected at 23 days post infection (DPI) with 10⁸ infected erythrocytes (iRBC) from another splenectomized calf and at 77 and 99 DPI with 10⁸ iRBC from culture. At day 127, the five cattle and three weight-matched, previously uninfected controls were infected with 10⁹ iRBC from culture. The packed cell volume (PCV) of all animals was monitored daily.

Immunofluorescence of Live Merozoites. Viable merozoites were collected as described above and reacted with the various antibodies by a previously described technique (McElwain et al. [1987], supra). The MoAb-containing ascites fluids were diluted 1/10 (40 ug/ml) in PBS. Immune bovine sera were diluted 1/10 in PBS. An equal volume of each antibody preparation was added to 100 ul of a merozoite suspension and incubated on ice for 1 hr. Each sample was centrifuged at 3,000 x g for 10 min, and the merozoites were washed twice in cold PBS. The samples were then suspended in the appropriate rhodamine-conjugated second antibody (1/40 dilution in PBS) (Kirkegaard and Perry, Inc., Gaithersburg, MD) and incubated on ice for 1 hr. After being washed, the merozoites were suspended in 6-CFDA and incubated for 20 min at room temperature. The merozoites were then centrifuged at 3,000 x g for 10 min at 4°C, suspended in 50 ul of PBS, and examined in wet mounts with appropriate filters for rhodamine (antibody binding) and fluorescein (6-CFDA viability) (546 to 590 nm and 450 to 520 nm, respectively).

Surface Radioiodination. Purity of the gradient separated merozoites was also examined by direct light microscopy of Giemsa stained smears and by transmission electron microscopy of selected samples fixed in 2% v/v glutaraldehyde

in 0.1 M potassium phosphate buffer containing 1% w/v sucrose. Parasites were often arranged in clumps and mixed with very rare erythrocyte ghosts (<0.1%). Merozoites were surface radioiodinated by a previously described lactoperoxidase catalyzed method (Palmer, G.H., and T.C. McGuire [1984] J. Immunol. 133:1010-1015).

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Donor erythrocytes (nRBC) from uninfected control cultures were collected, washed three times in PBS, and radiolabeled identically. An equivalent number of nRBC ghosts were prepared by lysing washed uninfected cells from control cultures by freeze/thaw in liquid nitrogen. Ghosts were washed free of hemoglobin in PBS by centrifugation at 35,000 x g, 20 min, 4°C and discarding the supernatant until it was clear. The final pellet was resuspended in PBS for radioiodination by lactoperoxidase.

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Metabolic Radiolabeling of Merozoites. Metabolically radiolabeled parasite proteins from calf-derived merozoites were prepared for use in immunoprecipitation experiments according to the methods of McElwain et al. (1987, supra) except that cultures containing 100 uCi of [35S]-methionine (35S-Met; New England Nuclear, Boston, MA) per 3 x 10° erythrocytes were incubated at 37°C for 8-9 hr in a Forma Scientific water jacketed incubator instead of a candle jar. Parasites cultivated in vitro were metabolically radiolabeled using normal growth medium (Goff and Yunker, 1986) or D,L-methionine-free medium, addition of 20-400 uCi/ml 35S-Met, 3H-myristic acid, or 3H-glucosamine, and incubation of cultures for 12-20 hr at 37°C. Erythrocytes containing radiolabeled merozoites were solubilized in lysis buffer, TCA-precipitable radioactive counts were determined by a filter paper technique (New England Nuclear), and samples were frozen at -70°C until used.

Phase Separation in TRITONTMX-114. Washed iRBC's from ³⁵S-methionine labeled cultures were lysed in 10 mM Tris, 154 mM NaCl pH 7.4, 1% (v/v) TRITONTMX-114, 1 mM phenylmethylsulfonyl fluoride (PMSF) at 0-4°C and frozen at -20°C. For protein separation, the antigen extract was first processed as described for immunoprecipitation. 10⁷ protein bound counts per minute (CPM) in a volume of 2.0 ml was laid over a 2.0 ml cushion of 6% (w/v) sucrose, 10 mM

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Tris, 154 mM NaCl, 1 mM PMSF in a 15 ml conical tube (Bordier, C. [1981] Exp. Parasitol. 20:125-129). The tube was incubated at 37°C for 5 min to allow clouding of the protein extract and then centrifuged at 750 x g at room temperature for 5 min in a swinging bucket rotor. The detergent phase was seen as a thick, oily 100-200 ul pellet and the overlying aqueous phase and sucrose cushion were each removed to separate tubes. The phase separation was repeated twice by adding 200 ul of 15% (v/v) TRITONTMX-100 in 10 mM Tris, 154 mM NaCl to the aqueous phase, dissolving the detergent on ice, and re-extracting at 37°C over the same sucrose cushion. The three detergent phases resulting from centrifugation were mixed with 10 mM Tris, 154 mM NaCl at 0-4°C, combined, and TCA-precipitable radioactivity counted along with the aqueous phase.

Immunoprecipitation. Immune sera were used either unadsorbed or adsorbed three times with an equal volume of packed intact nRBC's and three times with an equal volume of nRBC ghosts. Radiolabeled B. bovis or bacterial lysate was processed as described previously (Palmer and McGuire [1984], supra) and incubated overnight at 4°C with 15 ul of bovine serum or 15 ul of serum diluted in Veronal buffered saline (VBS) pH 7.4, 1% (v/v) Nonidet P-40 (NP-40). 150 ul of 10% (v/v) formalinized Protein G-bearing Streptococcus (Omnisorb, Calbiochem, San Diego, CA) in VBS pH 7.4, 1% (v/v) NP-40, 0.1% (w/v) gelatin was added and incubated for 2 hr at 4°C (Akerstrom, B., T. Brodin, K. Reis, and L. Bjorck [1985] J. Immunol. 135:2589-2592). The precipitates were washed twice with VBS, 1% (v/v) NP-40; four times with VBS, 2 M NaCl, 1% (v/v) NP-40, 10 mM ethylenediaminetetraacetic acid (EDTA); and twice more with VBS, 1% (v/v) NP-40. Alternatively, 5 ul of bovine serum, 10 ul of rabbit serum, or 5 ug monoclonal antibody were incubated for 30 minutes at 4°C with radiolabeled lysate. Rabbit anti-bovine or rabbit anti-mouse immunoglobulin sera were added and incubated for 30 minutes at 4°C, followed by 10% v/v protein-A-bearing Staphylococcal aureus for 30 minutes at 4°C. Immune complexes were washed seven times with TEN buffer (20 mM Tris-HCl, 5 mM EDTA, 0.1 mM NaCl, 15 mM NaN₃, pH 7.6) containing Nonidet P-40, and for the second through fifth washes, 2 M NaCl, by centrifuging at $1250 \, x$ g. The precipitated protein was eluted by a described method and either frozen at -20°C or loaded directly onto a polyacrylamide gel (Palmer and McGuire [1984], supra).

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Sodium Dodecyl Sulfate Polyacrylamide Gel Electrophoresis (SDS-PAGE) of Immunoprecipitates and Autoradiography. Immunoprecipitates were electrophoresed under reducing conditions in 7.5 to 17.5% continuous gradient polyacrylamide gels (Takac, B. [1979] In: Immunological Methods. T. Lefkovitz and B. Persin, eds., p. 81, Academic Press, New York). ¹⁴C-labeled protein standards used for molecular weight determination were myosin, 200 kDa; phosphorylase b, 92.5 kDa; bovine serum albumin, 69 kDa; ovalbumin, 46 kDa; carbonic anhydrase, 30 kDa; and lysozyme, 14.3 kDa (Amersham Corp., Arlington Heights, IL). For ¹²⁵I-labeled antigens, gels were fixed in 30% (v/v) methanol, 10% (v/v) acetic acid, vacuum dried, and exposed to Kodak XAR-2 X-ray film with an intensifying screen at -70°C. For ³H and ³⁵S-labeled immunoprecipitates, gels fixed in 30% (v/v) methanol, 10% (v/v) acetic acid, 10% (w/v) trichloroacetic acid (TCA) were impregnated with En³Hance (New England Nuclear Corp., Boston, MA) prior to drying and exposure to X-ray film at -70°C (Palmer and McGuire, 1984).

Immunoblotting. Immunoblotting of merozoite and recombinant lysogen proteins using monoclonal antibodies was performed as outlined (McElwain et al., 1987) using standard procedures (Towbin, H. and J. Gordon [1984] J. Immunol. Methods 72:313-340). Parasite antigen for immunoblotting was prepared from MASP culture flasks with approximately 25% parasitemia. Briefly iRBC's and nRBC controls were collected, washed two times in cold Puck's saline-G and two times in cold PBS, resuspended in PBS, counted, and frozen at -20°C. To remove hemoglobin from lysed cells, the samples were thawed and washed in cold PBS (43,000 x g, 20 min, 4°C) until the discarded supernatant was clear. The final pellet was extracted in lysis buffer, processed identically to radiolabeled antigen for immunoprecipitation, and aliquoted for freezing at -20°C. A volume corresponding to 2.5 x 10⁷ iRBC's or an equivalent total number of nRBC's (10⁸) was mixed with 3X SDS-PAGE sample buffer, boiled for 3 minutes, electrophoresed in a 7.5 to

17.5% continuous gradient polyacrylamide gel, and then electrophoretically transferred overnight to a nitrocellulose membrane (Towbin and Gordon, 1984). Immunoblotting using bovine antisera was performed as follows: The nitrocellulose was washed three times quickly in VBS pH 7.4, 0.25% (v/v) TWEENTM20, 0.25% (w/v) gelatin (blocking buffer), incubated 4-6 hr in blocking buffer, cut into strips, and each strip reacted overnight at room temperature with immune serum diluted in blocking buffer. The nitrocellulose strips were then washed three times in blocking buffer and two times in VBS 0.1% (w/v) gelatin prior to incubation for 2 hours at room temperature with ¹²⁵I-Protein G (Amersham Corp.) in VBS pH 7.4, 0.1% (w/v) gelatin (Akerstrom, B., T. Brodin, K. Reis, and L. Bjorck [1985] J. Immunol. 135:2589-2592). The strips were washed twice with VBS 0.1% (w/v) gelatin and four times with 1 M NaCl, 10 mM EDTA, 0.25% (v/v) TWEENTM20. They were then air-dried, taped to cardboard, and exposed to X-ray film with an intensifying screen at -70°C.

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Dot Blot Immunoassay. Because MoAbs were used to screen the lambdagt11 genomic library for clones expressing recombinant surface proteins, they were first evaluated for their ability to bind native antigen applied to nitrocellulose filters. 6-CFDA-positive merozoites were obtained from frozen blood stabilates, lysed in buffer containing 50 mM Tris, 5 mM EDTA, 5 mM iodoacetamide, 1 mM PMSF, 0.1 mM N-alpha-p-tosyl-L-lysyl chloromethyl ketone (TLCK) and 1% NP-40 (lysis buffer), and frozen at -70°C until use. Aliquots of 1 ul containing either 10⁷, 10⁶ or 10⁵ merozoites were spotted onto nitrocellulose filters and air dried. Corresponding numbers of similarly lysed noninfected bovine erythrocytes were spotted on for control. Nitrocellulose filters with spotted antigen were washed three times (10 min each) in buffer containing 10 mM Tris (pH 8.0), 150 mM NaCl, 0.05% TWEENTM20, and 0.1 mM PMSF (TNTP) then incubated in TNTP with 5% nonfat dry milk for 1 hr to block unbound sites. Filters were washed three times in TNTP plus 5% milk, incubated in the same buffer containing 2 ug/ml of specific surfacebinding MoAb (1/2 hr), washed three times, incubated for 1/2 hr in a 1:5000 dilution of rabbit anti-mouse immunoglobulin (prepared in our laboratory) in TNTP plus 5%

milk. After three washes, the filters were incubated for 1/2 hr in TNTP plus 5% milk containing 5 x 10⁶ CPM of ¹²⁵I-labeled Protein A, washed sequentially with TNTP, TNTP plus 0.1% TRITONTMX, and TNTP, then dried and examined by autoradiography.

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Following are examples which illustrate procedures, including the best mode, for practicing the invention. These examples should not be construed as limiting. All percentages are by weight and all solvent mixture proportions are by volume unless otherwise noted.

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Example 1 - Generation of Monoclonal Antibodies Against Surface Epitopes

Partially purified merozoites of B. bovis obtained from frozen blood stabilates were used to immunize BALB/c mice for hybridoma production. Each mouse received an initial subcutaneous immunization of 10⁷ 6-CFDA-positive organisms in Freund's complete adjuvant followed by 3-4 subcutaneous immunizations of 107 6-CFDA-positive organisms in Freund's incomplete adjuvant at 2-4 week intervals. Mouse serum was titered by IFA-live (Barbet, A.F. and T.C. McGuire [1978] Proc. Natl. Acad. Sci. USA 75:1989-1993) after the last immunization, and mice with high titers (≥1:1000) received an intravenous booster immunization of 10⁶ 6-CFDApositive organisms in sterile PBS. Three days later the mice were killed and their spleen cells fused to SP2/0 myeloma cells using standard procedures (McGuire, T.C., L.E. Perryman, and W.C. Davis [1983] Amer. J. Vet. Res. 44:1284-1288). Hybridoma supernates were screened first for convenience by IFA-fixed (Ross, J.P.J and K.F. Lohr [1968] Res. Vet. Sci. 9:557-562). Positive supernates were then screened by IFA-live using stabilate-derived merozoites in order to identify surface reactive MoAbs (McElwain et al. [1987]). The MoAbs were first screened on fixed infected erythrocyte preparations, and four MoAbs were selected for further evaluation because of their distinctive patterns of fluorescence. These patterns included staining of the merozoite cytoplasm and membrane (BABB35A4), merozoite membrane (BABB90C₄), and merozoite cytoplasm (BABB93A₁) and a 5

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single, punctate reaction appearing polar in location on merozoites (BABB75). The four MoAbs all retained their original specificities after the hybridoma cells were cloned twice and used to produce ascites fluids.

Example 2 - Immunoprecipitation of Surface Radioiodinated and Metabolically Radiolabeled Proteins

Spontaneously released merozoites for surface binding and labeling experiments were isolated on Percoll gradients. A large proportion retained their surface coat and >80% were viable, as determined by 6-CFDA staining. On three occasions, 10⁸ of these isolated merozoites were inoculated intravenously into susceptible, splenectomized calves. In each case, infection was achieved with a prepatent period similar to that in calves that received an equivalent number of infected erythrocytes from a blood stabilate. Also, in vitro cultures have been routinely reestablished after introduction of these isolated merozoites.

Radioiodinated merozoite preparations were immunoprecipitated with the MoAbs described above to confirm the outer surface or cytoplasmic location of the reactive epitopes. BABB35A₄ precipitated a major protein of 42 kDa and a minor protein of 37 kDa. BABB75 and BABB90C₄ precipitated single proteins of 60 and 85 kDa, respectively.

To determine which parasite proteins were recognized by the bovine immune system, we used twofold serial dilutions of immune bovine sera to immunoprecipitate metabolically labeled preparations. Proteins with relative molecular masses ranging from approximately 16 to >200 kDa were recognized by the immune bovine sera. Among the proteins recognized were those identical in molecular mass to those precipitated by BABB35A₄ (42 and 37 kDa), BABB75 (60 kDa), and BABB93A₁ (145 kDa). In addition, proteins of 145, 42, 120 and 75 kDa appeared to be immunodominant, as they were precipitated by immune bovine sera at the greatest dilution tested.

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Example 3 - Further Identification of Merozoite Surface Proteins

Subinoculation of 5 x 10^7 infected erythrocytes (iRBC) of a cloned <u>B. bovis</u> line from a splenectomized calf into 4-5 month old cattle caused a 39% reduction in packed cell volume (PCV) (range 31-47). Calves were re-infected three times with approximately 108 iRBC of the cloned isolate and then challenged with 109 iRBC in concert with three previously uninfected control animals. Only the initial infection caused a significant reduction in PCV when compared to the PCV during the week prior to each infection. Control cattle in the final challenge experiment experienced a 28% reduction in PCV (p < or = 0.0005 when compared to previously infected cattle; Student's paired t test).

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Merozoites spontaneously released from culture and purified on Percoll gradients were 95-100% viable by 6-CFDA staining. In all five animals, immunoprecipitation of surface radioiodinated proteins with immune sera that had been extensively adsorbed against donor erythrocytes (nRBC) and erythrocyte ghosts identified seven dominant surface proteins with relative molecular weights of 250, 125, 98, 85, 55, 42, and 37 kilodaltons. The 250 kDa protein does not enter the resolving gel in a standard 14 cm 7.5-17.5% polyacrylamide gel but is clearly resolved in a 25 cm gel. An eighth protein of 25 kDa is immunoprecipitated by immune sera from two calves. Control immunoprecipitation of identically radioiodinated intact nRBC and nRBC ghosts revealed no specific bands on SDS-PAGE.

Adsorbed immune sera was used to immunoprecipitate 35S-methionine metabolically labeled parasite proteins which were immunoprecipitated surface-iodinated merozoite proteins in a polyacrylamide gel. The immunoprecipitable ³⁵S antigen profile is identical in all five protected animals. The 125, 98, 85, 55, 42, and 37 kDa antigens comigrate perfectly with metabolically The 25 kDa surface protein that is not identified by labeled proteins. immunoprecipitation of methionine labeled antigen does comigrate with a glycoprotein that is metabolically labeled with ³H-glucosamine. An ³⁵S-methionine labeled 25 kDa protein can be precipitated from other 35S-antigen preparations.

Example 4 - Immunogenicity

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While the reactivity of immune sera against the majority of ³⁵S-labeled proteins can be diluted out, sequential serum dilutions (1:160 - 1:640) selectively precipitate the 125, 55, and 42 kDa proteins that were also surface labeled. Because this method is dependent on the specific radioactivity of labeled proteins, dilute sera was also examined for its ability to react with parasite antigens by immunoblotting. Compared to undiluted serum, immune sera diluted 1:500 recognizes a limited number of blood stage proteins including the 125, 85, 55, and 42 kDa surface antigens. The 42 kDa protein is consistently recognized even at dilutions of greater than or equal to 1:16,000.

Example 5 - Further Characterization of Proteins

The immunodominant 42 kDa merozoite surface protein was further characterized as an integral membrane protein based on its hydrophobic nature in phase separated TRITONTMX-114 solution. By definition, integral membrane proteins have a hydrophobic domain that allows interaction with the hydrophobic core of the lipid bilayer and with non-ionic detergents. Parasite proteins were metabolically labeled in culture with ³⁵S-methionine and solubilized in 1% TRITONTMX-114 at 0-4°C. The antigen preparation was warmed above the detergent's cloud point (20°C) and separated into aqueous and detergent phases by centrifugation. Immunoprecipitation from each phase and the starting solution shows that the 42 kDa antigen partitions into the detergent phase.

In order to better characterize merozoite surface antigens, the parasite was examined for the ability to incorporate ³H-glucosamine and ³H-myristic acid into immunoprecipitable proteins. Comigration on a polyacrylamide gel shows that three of the surface labeled proteins (55, 42, and 25 kDa) are glycosylated and the 42 kDa glycoprotein is myristylated.

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Example 6 - Further Studies on Immunogenicity

Antiserum C151, which was used for immunoprecipitations, was collected from a spleen-intact cow 60 days after experimental infection with a cryopreserved Mexico isolate blood stabilate of <u>B. bovis. B. bovis</u> proteins were metabolically labeled in microaerophilus stationary-phase culture by incubation in methionine-deficient medium for 18 to 24 hr with 10 uCi of [35S]methionine per ml. Antiserum C151 immunoprecipitated homologous Mexico isolate proteins biosynthetically labeled with [35S]methionine with molecular weights ranging from 14,500 to greater than 200,000. Serial dilution of this antiserum resulted in a decrease in the number of proteins recognized. Proteins reactive with serum diluted 1:40 had relative molecular weights of 145,000, 120,000 (doublet), and 42,000, while the 42,000 molecular weight protein was still recognized by serum diluted 1:80.

Example 7 - B. bovis Proteins with Isolate-Common Epitopes

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Five different antisera obtained from cattle after recovery from acute infection with <u>B. bovis</u> in Honduras were able to immunoprecipitate most of the Mexico isolate <u>B. bovis</u> proteins precipitated by C151 antiserum. The 120,000 and 42,000 molecular weight proteins recognized by 1:40 dilutions of C151 antiserum were also recognized by 1:25 dilutions of the Honduran antisera.

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Example 8 - B. bovis Proteins with Species-Common Epitopes

Antiserum B85 was collected from a spleen-intact calf 25 days after experimental infection with a cryopreserved Mexico isolate of <u>B. bigemina</u>. This antiserum, which had an indirect fluorescent-antibody titer of 1:1,600 against the Mexico <u>B. bigemina</u> isolate, reacted with the Mexico <u>B. bovis</u> isolate at a titer of 1:64. Antiserum C151 (anti-<u>B. bovis</u> Mexico isolate) had indirect fluorescent-antibody titers of 1:5,120 and 1:640 against <u>B. bovis</u> and <u>B. bigemina</u>, respectively. Antiserum B85 immunoprecipitated eight [35S]methionine-radiolabled proteins of <u>B. bovis</u>. Four of the eight <u>B. bovis</u> proteins immunoprecipitated by B85 antiserum (120,000, 59,000, 53,000, and 19,000 molecular weight) also had isolate-common

epitopes. In addition, the 120,000 molecular weight protein was one of the proteins recognized by C151 serum antibodies diluted 1:40.

Example 9 - Identification of Monoclonal Antibodies

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Using similar techniques, additional MoAbs specific for surface-exposed epitopes on live merozoites were identified. All of the identified MoAbs are listed in Table 1. The MoAbs reacted with the outer surface of culture- or stabilatederived merozoites in either a punctate (restricted to a discrete region on the merozoite surface) or a homogenous (over the entire surface of the merozoite) pattern when examined by IFA-live.

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Table 1. Monoclonal antibodies generated against surface proteins on merozoites of Babesia bovis.

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MoAb	Isotype	MW of Reactive Protein (x 10 ⁻³ kd)
14102 10	isotype	(x 10 - kd)
23.8.34.24	G_3	22 5
BABB75	G_{2b}	60
MBOC79	G_1	60
23.53.156	G_{2a}	60
23.38.120.8	G_1	60
23.70.174.83	G_1	44
BABB35A₄	G_{2a}	42
23.3.16	G_1^{-}	42
23.10.36	G_{2b}	42
23.28.57.108	G_{2a}^{2a}	16
BABB90C ₄	G_1^{ra}	85
BABB93A	G_{2a}	145

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All MoAbs reacted with merozoite antigen in dot immunoassay and allowed detection of specific surface-exposed determinants in preparations of 105 lysed

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merozoites. The parasite specificity of these MoAb-reactive determinants was confirmed by immunoprecipitation of metabolically-radiolabeled parasite proteins of M_r 16 kDa, 42 kDa, 44 kDa, 60 kDa, and 225 kDa. These proteins have been designated Bo16, Bo42, Bo44, Bo60, and Bo225, respectively. Bo225 was routinely visualized as a tightly spaced doublet when immunoprecipitated with MoAb 23.8.34.24.

Seven monoclonal antibodies and the Mexico isolate of <u>B. bovis</u> have been deposited with the American Type Culture Collection (ATCC), 12301 Parklawn Drive, Rockville, Maryland 20852 USA. The cultures have been assigned the following accession numbers by the repository:

	Biological Material	Deposit number	Deposit date
	MoAb 23.38.120.8	HB 10111	May 2, 1989
	MoAb BABB93A ₁	HB 10112	May 2, 1989
15	MoAb 23.8.34.24	HB 10113	May 2, 1989
	MoAb 23.70.174.83	HB 10114	May 2, 1989
	MoAb BABB35A₄	HB 10115	May 2, 1989
	MoAb 23.28.57.108	HB 10377	March 7, 1990
	MoAb BABB90C₄	HB10117	May 2, 1989
20	Babesia bovis, Mexico Isolate	ATCC 40601	May 3, 1989
		- 	14145 3, 1709

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The subject cultures have been deposited under conditions that assure that access to the cultures will be available during the pendency of this patent application to one determined by the Commissioner of Patents and Trademarks to be entitled thereto under 37 CFR 1.14 and 35 USC 122. The deposits are available as required by foreign patent laws in countries wherein counterparts of the subject application, or its progeny, are filed. However, it should be understood that the availability of a deposit does not constitute a license to practice the subject invention in derogation of patent rights granted by governmental action.

Further, the subject culture deposits will be stored and made available to the public in accord with the provisions of the Budapest Treaty for the Deposit of Microorganisms, i.e., they will be stored with all the care necessary to keep them viable and uncontaminated for a period of at least five years after the most recent request for the furnishing of a sample of the deposit, and in any case, for a period of at least 30 (thirty) years after the date of deposit or for the enforceable life of any patent which may issue disclosing the cultures. The depositor acknowledges the duty to replace the deposits should the depository be unable to furnish a sample when requested, due to the condition of the deposit(s). All restrictions on the availability to the public of the subject culture deposits will be irrevocably removed upon the granting of a patent disclosing them.

Example 10 - Construction of a Lambda-gt11 Expression Library

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Partially purified and viable merozoites free from contaminating bovine leukocytes were obtained from frozen stabilates. Merozoites (1.4 x 108 6-CFDApositive) were lysed in 10 mM Tris, 1 mM EDTA (TE buffer, pH 7.4) containing 2% SDS, and the suspension was treated with DNAse-free RNAse A (100 ug/ml) followed by Proteinase K (100 ug/ml). Genomic DNA was isolated from the suspension by sequential phenol, phenol/chloroform, chloroform, and ether extractions followed by ethanol precipitation at 0-7°C in the presence of 2 M ammonium acetate. The DNA pellet was washed once with 70% ethanol, lyophilized, resuspended in TE buffer and stored at 4°C. The concentration and purity of the DNA were assessed by spectrophotometry and agarose gel electrophoresis. The DNA was sheared into fragments of between 4-8 kb by repeated passages through a 25 gauge hypodermic needle (Young, R.A., B.R. Bloom, C.M. Grosskinsky, J. Ivanyi, D. Thomas, and R.W. Davis [1985] Proc. Natl. Acad. Sci. USA 82:2583-2587) and the fragments prepared for ligation into the EcoRI site of the lambda-gt11 expression vector (Promega Biotec, Madison, WI).

First, fragments were methylated with <u>Eco</u>RI methylase (Promega Biotec) and blunt-ended using the large fragment of <u>E. coli</u> DNA polymerase I (Klenow

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Fragment, Bethesda Research Laboratories, Gaithersburg, MD). <u>Eco</u>RI linkers (BRL, Gaithersburg, MD), end-labeled with ³²P transferred from 5'-[gamma-³²P]ATP in a kinase reaction (Huyhn et al., 1985), were ligated to fragment termini using T4 DNA ligase (Bethesda Research Laboratories). Free linkers were separated from fragments with <u>Eco</u>RI termini by size fractionation on a Sephacryl S-400 column (Pharmacia AB, Uppsala, Sweden) following digestion of the reaction mixture with <u>Eco</u>RI endonuclease.

Fractions containing fragments with EcoRI cohesive termini but free of nonligated linkers were pooled, butanol-extracted to reduce volume, extracted twice with ether, ethanol-precipitated, lyophilized, and resuspended in TE buffer. Fragments were then ligated into the EcoRI site of the lambda-gt11 expression vector which resulted in the insertion of parasite DNA into the \(\beta\)-galactosidase structural gene (lacZ) of the bacteriophage (Young, R.A. and R.W. Davis [1983] Proc. Natl. Acad. Sci. USA 80:1194-1198). Ligated DNA was packaged into gamma phage heads (Gigapack Gold Packaging Extract, Stratagene Cloning Systems, San Diego, CA) and the resultant library was amplified in E. coli strain Y1090 as described previously (Young and Davis, 1983). The amplified library was stored in sterile SM buffer (0.1 M NaCl, 8 mM MgSO₄·7H₂O, 50 mM Tris [pH 7.5], 2% gelatin) at 4°C.

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Example 11 - Identification of Recombinant Phage Expressing Parasite Surface-Exposed Proteins

Recombinant phage expressing proteins with surface-exposed epitopes were identified by immunoscreening plaques with MoAbs. Enough recombinant phage to give 10⁵ plaque forming units (pfu)/150 mm diameter petri dish were used to infect E. coli host Y1090 by incubation at 37°C for 20 min in LB medium. Infected cells were added to LB top agar (55°C) containing 100 ug/ml ampicillin and 10 mM MgCl₂ and plated out on 150 mm diameter LB agar plates. Plates were incubated at 42°C for 4 hr to allow plaque formation without concomitant expression of fusion protein. LacZ-directed gene expression was then switched on by overlaying each

plate with a dry nitrocellulose filter saturated previously with 10 mM IPTG and incubating the plates at 37°C for 8-10 hr. After incubation, nitrocellulose filters with bound proteins were marked, removed from the plates, and processed as described previously for dot blot immunoassay. Single plaques expressing recombinant surface epitopes of interest were identified by autoradiography, picked from plates, and rescreened and picked three more times to insure purity of the recombinant phage, stability of the DNA insert, and reliability of recombinant protein expression. Other hosts, such as Salmonella, can be transformed by suitable procedures well known to those in the art.

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Approximately 4.2 x 10⁶ recombinant plaques were screened with MoAbs listed in Table 1. Two recombinant clones (lambda-Bo44-15, lambda-Bo44-16) were identified that express a recombinant protein that reacts with MoAb 23.70.174.83 (anti-Bo44) and two (lambda-Bo220-1 and lambda-Bo220-2) that express a recombinant protein that reacts with MoAb 23.8.34.24 (anti-Bo225). When lambda-Bo44-15 and lambda-Bo44-16 were digested with EcoRI, inserts (Bo44-15, Bo44-16) of 1.25 kb were visualized for each recombinant clone. IPTG-induced lysogen preparations of lambda-Bo44-15 consistently produced stronger signals in dot blot immunoassay than did lambda-Bo44-16, and for this reason, lambda-Bo44-15 was chosen for further analysis and use in immunization trials.

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Example 12 - Induction of Recombinant Proteins with IPTG

E. coli host strain Y1089 was lysogenized with lambda-gt11 (control) and each of the recombinant clones using standard procedures (Huyhn, T.V., R.A. Young, and R.W. Davis [1985] "Constructing and Screening cDNA Libraries in lambda-gt10 and lambda-gt11," In: DNA Cloning, Vol. 1: A Practical Approach [Glover, D.M., ed.], pp. 49-78, IRL Press, Washington, D.C.). Lysogenized bacteria were examined by dot blot immunoassay in order to determine the ability of clones to produce recombinant protein after induction with IPTG. Each of the recombinant clones, lambda-gt11-infected \underline{E} . coli Y1089, and noninfected Y1089 controls were grown at room temperature to OD_{600} =0.8-1.2. At this time, an

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aliquot was removed, centrifuged, and lysed with lysis buffer (1% NP-40). The remaining cells were heated rapidly to 42-45°C for 20 min, IPTG was added to 10 mM, and the cells were incubated in a shaking incubator at 37°C for 1-2 hr to induce protein expression. After incubation, cell cultures were adjusted by addition of LB medium to their OD₆₀₀ prior to addition of IPTG. At this time, an aliquot was removed, centrifuged, and the pellet lysed in lysis buffer. Three ul aliquots of pre- and post-induced bacterial lysates were spotted onto nitrocellulose in triplicate and probed with MoAbs specific for surface proteins or an irrelevant MoAb control (CAEV 4Al) in dot blot immunoassays. Crude lysates of bacteria producing recombinant protein were prepared and stored at -20°C or -70°C until use.

Dot blot and Western blot analysis of lysates of bacteria lysogenized with lambda-Bo44-15 confirmed the inducibility of rBo44-15 with IPTG and its expression as a β -galactosidase fusion protein. rBo44-15 was visible as a doublet of M_r 150 kDa and 165 kDa in Western blots of IPTG-induced preparations probed with MoAb 23.70.174.83. In contrast, rBo44-15 was visible as a single band (165 kDa) in Western blots of identical antigen preparations probed with anti- β -galactosidase. Western blots of lambda-Bo44-15 lysogen preparations probed with an irrelevant IgG₁ control MoAb (5.90.1) showed no reactivity, thus confirming the specificity of reaction observed with MoAb 23.70.174.83.

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Example 13 - Purification of Recombinant Proteins

MoAb 23.70.174.83 was purified from ascitic fluid by ammonium sulfate precipitation and DEAE cellulose chromatography and then coupled to Sepharose 4B for immunoaffinity purification of rBo44-15. Solubilized and sonicated rBo44-15 lysogen preparations were applied to the affinity column, the column was washed repeatedly, then adherent recombinant protein was eluted with 0.1 M diethylamine (pH 11.5) containing 0.5% deoxycholate. Elutes were collected directly into 1 M Tris (pH 8.5) then dialyzed against PBS to remove detergents. Aliquots of partially purified protein preparations were boiled for 10 min in SDS sample buffer, subjected to SDS-PAGE, silver stained and examined by Western blot immunoassay

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to verify the presence and purity of recombinant protein. Total protein concentration of the preparations was determined using a bicinchoninic acid protein assay (Pierce Chemical Co.).

Partially purified rBo44-15 from affinity column chromatography contained several high M_r proteins ranging from approximately 94 Kd to >165 Kd as well as several lower M_r proteins ranging from 26 Kd to 50 Kd. Western blot analysis of column-purified protein preparations revealed two major bands of reactivity at M_r 165 Kd and 150 Kd that correspond to two major bands present in silver-stained gels. In addition, several bands of lower M_r (26-31 Kd) were observed in Western blots of affinity-purified recombinant protein that were not observed in Western blots of solubilized lysogen preparations prior to affinity purification. These data indicate that immunoaffinity chromatography results in partial degradation of the recombinant molecule without a concomitant loss of integrity of the MoAb-reactive epitope.

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Example 14 - Confirmation of the Presence of Surface-Exposed Epitopes on Recombinant Molecules

Five Holstein-Freisian bull calves were each immunized intramuscularly (i.m.) with 100-125 ug of affinity column-purified recombinant protein in Freund's complete adjuvant, followed by four to five additional immunizations at three week intervals of recombinant protein in Freund's incomplete adjuvant. Control calves were immunized similarly with 100 ug ovalbumin. Within one week after the last immunization, calves were bled and their sera heat-inactivated and examined by IFA-live (Goff et al., 1988) to confirm the presence of antibodies to surface-exposed epitopes on merozoites. Hyperimmune bovine serum (KLK C151) and preimmune sera were used as positive and negative controls, respectively. Sera from calves immunized with either rBo44-15 or ovalbumin were used to immunoprecipitate metabolically radiolabeled merozoited proteins (McElwain et al., 1987) in order to verify the specificity of the antibody response.

Antibody titers in serum from calves immunized with partially purified rBo44-15 varied from 10⁻² to 10⁻⁵ as evidenced by IFA-live. In contrast, preimmune serum (B452-pre) and serum from ovalbumin-immunized calves showed reactivity with live merozoites at dilutions of 10⁻¹ and 10⁻², respectively. Antibody in all nondiluted sera or in low serum dilutions (10⁻¹, 10⁻²) bound to erythrocyte ghosts (infected and noninfected) as well as merozoites within ghosts.

Example 15 - Methods and Materials for Construction of DNA Probe

(a) Parasites and DNA Isolation

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Strains of parasites used in this study include a Mexico (M) and Australia (S strain) isolate of <u>B. bovis</u> and a Mexico isolate of <u>B. bigemina</u>. <u>Babesia bovis</u> (M) DNA for both the genomic library preparation and analysis of clones was derived from infected bovine erythrocyte cultures washed three times in phosphate buffered saline (PBS), pH 7.2, followed each time by centrifugation at 400xg. Babesia bovis (S) and B. bigemina DNA was similarly derived from infected calf blood depleted of buffy coats by three washes in PBS. Infected erythrocytes for isolates were differentially lysed in nine volumes of 0.42% NaCl, infected ghosts were pelleted at 400xg, lysed in 5 volumes of 10 mM Tris-HCl (pH 7.5), 10 mM ethylenediaminetetraacetic acid (EDTA), 100 mM NaCl, and 1% sodium dodecyl sulfate (SDS), incubated 16 hr with proteinase K (100 ug/ml), extracted with phenol:chloroform:isoamyl alcohol (24:24:1). DNA in the aqueous phase was spooled after addition of 2 volumes of cold ethanol, spooled DNA was dried and resuspended in 10 mM Tris-HCl (pH 7.4) and 1 mM EDTA (TE), treated with RNases A and Tl (15 ug/ml, 15 units/ml, respectively). The solution was reextracted, spooled, dried, and resuspended in TE for use.

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To obtain bovine leukocyte DNA, cells in the buffy coat of uninfected blood were processed similar to infected erythrocytes.

(b) Identification and Isolation of Recombinant B. bovis DNA

The preparation of the genomic library has been described. Briefly, <u>B. bovis</u> genomic DNA was sheared through a 26 gauge needle to sizes ranging from ç4-8

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Kb, methylated with <u>EcoRI</u> methylase, <u>EcoRI</u> linkers were added, DNA restricted with <u>EcoRI</u> and separated from digested linkers on sephacryl S-400 (Pharmacia), and DNA ligated into <u>EcoRI</u> digested and dephosphorylated arms of lambda-gt11. Recombinant phage were amplified in <u>Escherichia coli</u> strain Y1089 (Stratagene).

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DNA from phage plaques was adsorbed onto nitrocellulose and replicate filters were differentially hybridized to nick translated DNA (2X 10⁶ cpm/ml) from either <u>B. bovis</u> (M) or <u>B. bigemina</u> in 6X SSPE (1X SSPE:150 mM NaCl, 10 mM NaH₂PO₄, 1 mM EDTA, pH 7.4), 100 ug/ml denatured salmon sperm DNA, and 1% SDS at 65°C for 16 hr. Filters were washed twice in 2X SSPE and 1% SDS at room temperature for 20 min, and twice in 0.1X SSPE and 0.1% SDS at 65°C for 20 min. Dried filters were autoradiographed at -70°C. Recombinant phage hybridizing to <u>B. bovis</u> (M) but not detectably to <u>B. bigemina</u> DNA were purified through 3 rounds of rescreening and isolated for further analysis. Based on characteristics of hybridization to <u>B. bovis</u> (M) genomic DNA, inserts from some recombinants were cloned into plasmid pBS⁺ (Stratagene) to facilitate their analysis.

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(c) Southern and Dot Blot Assays

To investigate the genomic organization of candidate probe sequences, restriction fragments were separated electrophoretically on 0.7% agarose gels and transferred to nylon filters. Filters were then hybridized, as described above, in the presence of 10% dextran sulfate, to lambda-gt11 recombinant DNA or preparatively isolated insert DNA that was radioactively labeled. Final wash stringency was either 65°C or 50°C, as indicated.

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For dot blot analysis, DNA extracted as described above was spotted onto nylon membranes. Membranes were dried at room temperature, saturated with 0.5 M NaOH and 1.5 M NaCl, neutralized in 1 M ammonium acetate and 0.02 M NaOH, rinsed in 6X SSPE, and vacuum-dried at 80°C for 1 hr. Hybridization conditions were similar to those used for Southern blots. Sensitivity of probe sequences was determined for autoradiograms exposed to hybridization filters for approximately 16 hr at -70°C using an intensifying screen.

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Example 16 - Candidate DNA Probes for Detecting Babesia bovis in Infected Ticks and Cattle

DNA-DNA hybridization assays (DNA probes) are based on the fact that single-stranded DNA will reanneal only with a complementary strand of DNA whose sequence is homologous. DNA probes have been used as a means of detecting various infectious agents, and some are now used routinely in clinical microbiology laboratories. The identification of DNA sequences of <u>Babesia</u> spp. makes it possible to create DNA probes for the identification of these species. Therefore, one application of the identification and isolation of genomic sequences which encode babesial antigens is the use of the DNA fragments as DNA probes.

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The lambda-gt11 genomic DNA library of <u>Babesia bovis</u> was screened to identify DNA probe candidates for direct detection of the parasite in blood or ticks infected with the parasite. Two sequences (lambda-Bo6 and lambda-Bo25) demonstrated superior sensitivity and were analyzed in more detail. The insert size of lambda-Bo6 is 2.75 kilobase pairs (kb). An accurate estimate of the lambda-Bo25 insert was not possible since one of the <u>EcoRI</u> insert sites was lost during cloning. However, digestion of lambda-Bo25 with <u>EcoRI</u> and <u>KpnI</u> produces a fragment of 2.2 kb which is specific to this clone compared to wild type lambda-gt11. Since the <u>KpnI</u> site in lambda-gt11 occurs approximately 1 kb from the <u>EcoRI</u> cloning site, a minimum estimated size of the insert is 1.2 kb.

Inserts from the lambda clones were excised with <u>EcoRI</u> (lambda-Bo6) or <u>SstI</u> and <u>KpnI</u> (lambda-Bo25) and cloned into the plasmid BS⁺ (Stratagene), producing the two clones pBo6 and pBo25. Insert fragments preparatively isolated from these plasmid clones were radioactively labeled and used in dot and Southern blot analyses.

Neither sequence hybridized detectably to bovine DNA. pBo6 detected 100 pg of both a Mexico and an Australia isolate of <u>B. bovis</u>, but pBo6 also detects 1.0 ng of <u>B. bigemina</u> DNA under identical conditions. A unique characteristic of pBo6 is that it hybridizes to a 7.4 kilobase band in uncut genomic DNA of both <u>B. bovis</u> and <u>B. bigemina</u>. Similarity of restriction enzyme patterns of the pBo6 sequence in

genomic DNA from both geographic isolates suggests that this sequence is well conserved between geographic isolates of <u>B. bovis</u>. Thus, this sequence is a candidate DNA probe for detecting <u>B. bovis</u> infections in cattle and ticks.

pBo25 exhibited no detectable hybridization to bovine or <u>B. bigemina</u> DNA. This sequence detected 100 pg of homologous Mexico isolate DNA, but under identical conditions the sensitivity was reduced to 1 ng for Australia isolate DNA. Restriction enzyme analysis of the pBo25 sequence showed major differences in the number, size, and intensity of bands between the two <u>B. bovis</u> geographic isolates tested. Thus, this sequence can distinguish geographic isolates of <u>B. bovis</u>.

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Example 17 - Labeling of DNA Probe

In order to facilitate detection, DNA probes can be labeled in a variety of ways. For example, for biotin labelling, the DNA fragment preparation is adjusted to a concentration of 1 mg/ml (TE) and is mixed with photo-activatable biotin (PAB) at a ratio of 1:3 (DNA:PAB) in a 1.5 ml Eppendorf tube. The tube is placed in an ice bath 10 cm below a 275 W (GE RSM) sunlamp and the DNA + PAB is irradiated for 15 min. The DNA solution is then mixed with an equal volume of 0.1 M Tris-Cl (pH 9.0) and the volume adjusted to 100 ul with H₂0. The unincorporated PAB is extracted from the DNA by the addition of an equal volume of 2-butanol, vortexing, centrifuging briefly, and withdrawing the lower aqueous phase with a Pipetman. The extraction can be repeated to remove any traces of unbound PAB. 3 M NaOAc (pH 5.6) is added to the DNA solution to a final concentration of 0.3 M and the labeled DNA is precipitated by the addition of 3 volumes of ethanol.

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After the sample is cooled at -70°C for 15 min, the precipitated DNA is recovered by centrifugation for 10 min. The DNA pellet is dissolved in 10 mM Tris (pH 7.9) and 0.1 mM EDTA. The labeled probe DNA remains stable for 1 year if stored at -20°C.

A non-radioactive method of labeling the DNA probes may be desirable because: 1) the photoactivatable reactions are simple and rapid, 2) the sensitivity

is as high as ³²P-labeled probes, 3) the PAB-labeled probes have a long storage life, 4) these probes are relatively inexpensive, and 5) detection of bound probes is by simple colorimetric methods. The radioactive labeling of probes requires the use of 32P, which has a very short half-life (14 days) and is thus unstable and expensive. The use of radioactive probes would be limited because of cost, the dangers of radioactivity, strict requirements for disposal, and the need for licensing.

However, if for some reason the biotin-HRP method of labeling is not acceptable, the DNA fragments can be labeled with [gamma-P] 32 deoxy CTP by standard nick translation methods.

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Example 18 - Description of Recombinant DNA Sequence from B. bovis that Encodes an Immunoreactive Epitope Located on the Surface of Merozoites

The cloned insert DNA was excised from the lambda-gt11 vector and recloned into the plasmid BS⁺ (Stratagene), producing the clone pBo44-15. DNA templates for sequencing the insert of pBo44-15 were obtained by creating deletion libraries of this clone using exonuclease III and mungbean nuclease. A different deletion library was obtained starting at each end of the clone, which allowed sequencing of both strands of the insert DNA. The DNA sequence was obtained using the Sanger dideoxy method.

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The sequence of Bo44-15 insert DNA is shown in Figure 1. The insert is 1235 base pairs long. The amino acid sequence shown represents the one long open reading frame identified in the sequence. The open reading frame begins at position 1 and encodes a stop codon (TAA) beginning at position 568. This reading frame is in correct register for expression as a fusion protein of \(\beta\)-galactosidase in lambda-gt11, provided the clone is in the correct orientation. A notable feature of this open reading frame is that it would encode a 24 amino acid sequence beginning at AA position 85 which is tandemly repeated beginning at AA position 109. Comparison of these two putative repeats shows only two positions that differ between the repeats as shown below:

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PQRPAETQQTQDSAAPSTPAAPSP 108PQRPAETQQTQDSTAPGTPAAPSP 132

Numbers represent the beginning and ending amino acid position in the open reading frame for each repeat. Letters are the single letter code for amino acids. Asterisks below the aligned repeats indicate amino acid differences between the two repeats.

The potential significance of the repeat amino acid sequence is that such repeats are often immunodominant epitopes in surface proteins from a variety of other protozoan parasites, and they induce antibodies that protect against the diseases.

The DNA sequences coding for two other of the <u>B. bovis</u> proteins have also been discovered. The DNA sequence for the <u>B. bovis</u> surface proteins of 42 kDA and 60 kDA are shown in Figures 2 and 3, respectively. These sequences, or portions of the sequences, can be used as DNA probes as described in Examples 16 and 17. Also, the proteins produced from cells transformed with these sequences, or portions of these sequences, can be used for vaccines or in the preparation of monoclonal antibodies as described in the examples which follow.

The procedure for obtaining these sequences are described below:

B. bovis cDNA Expression Library. Erythrocytes from asynchronous B. bovis-infected blood cultures were washed three times in Puck's saline G and stored frozen in liquid nitrogen. Cells were thawed in lysis buffer containing 0.2 M NaCl, 0.2 M Tris-HCl pH 7.5, 1.5 mM MgCl₂, 2% SDS (w/v), and 200 μg/ml Proteinase K and then incubated in lysis buffer at 46°C for two hours (Bradley, J.E., G.A. Bishop, T. St. John, and J.A. Frelinger [1988] Biotechniques 6:114-116). The NaCl concentration of the lysate was adjusted to 0.5 M and poly [A⁺] RNA was isolated by batch adsorption with oligo(dT) cellulose (Bradley et al., supra). DNA eluted with 0.01 M Tris-HCl pH 7.5 was used to prepare a blood stage cDNA library in Lambda ZAP II (Stratagene, La Jolla, CA, USA) (Short, J.M., J.M. Fernandez, J.A.

Sorge, and W.D. Huse [1988] Nucl. Acids Res. 16:7583-7600) by a modified Gubler and Hoffman method using EcoRI adapters (Pharmacia LKB, Piscataway, NJ, USA) (Gubler, U., and B. Hoffman [1983] Gene 25:263-269). The cloned insert in plaque purified lambda phage was subcloned into Bluescript SK(-) phagemid using the in vivo excision capabilities of Lambda ZAP II (Short et al., supra).

Immunoscreening. Plaque lifts ont isopropyl thiogalacto-pyranoside soaked nitrocellulose were screened using monospecific rabbit anti-Bv42 antisera (R-914) followed by 125I-Protein A and autoradiography, or with anti-Bv60 monoclonal antibody (23.38.120.8) followed by rabbit anti-murine immunoglobulin, ¹²⁵I-Protein A and autoradiography (Young, R.A., and R.w. Davis [1983] Proc. Natl. Acad. Sci. USA 80:1194-1198; Reducker, D.W., D.P. Jasmer, W.L. Goff, L.E. Perryman, W.C. Davis, and T.C. McGuire [1989] Mol. Biochem. Parasitol. 35:239-248). Rabbit R-914 had been immunized with native Bv42 protein immunoaffinity purified using BABB35A₄, a previously described monoclonal antibody (Goff, W.L., W.C. Davis, G.H. Palmer, T.F. McElwain, W.C. Johnson, J.F. Bailey, and T.C. McGuire [1988] Infect. Immun. 56:2363-2368). Positive Bv42 plaques were tested for reactivity with monoclonal antibodies that recognize a Bv42 surface exposed epitope (Goff et al., supra; Reducker et al., supra) as well as an isotype control monoclonal antibody and normal rabbit serum. Recombinant phagemid excised from positive, plaque purified lambda phage was tested for expression by a similar method using colony lifts from transformed, ampicillin resistant E. coli (XL1-Blue strain) (Young and Davis, supra).

Restriction Enzyme Digestion. Lambda rBv42 phagemid DNA was isolated from bacteria by anion exchange chromatography (Qiagen Inc., Studio City, CA) and restriction enzyme digested by standard methods (Maniatis, T., E.F. Fritsch, and J. Sambrook [1982] Molecular Cloning: A Laboratory Manual. Cold Spring Harbor Laboratories, Cold Spring Harbor, New York, 545 pp.).

Example 19 - Vaccines

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Vaccines may be produced from the polypeptides expressed by the parasites themselves or by cells which have been transformed with DNA fragments from

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<u>Babesia</u>. By introducing these polypeptides, along with a pharmacologically suitable vehicle or adjuvant, into the animal host, that host can be induced to generate immunological protection against <u>Babesia</u>. The preparation of such a vaccine composition is within the skill of one trained in the medical and immunological sciences. Vaccines may utilize entire polypeptides or epitopes with immunological activity.

Example 20 - Monoclonal Antibodies

Appropriate mice can be immunized with antigens of, or cells expressing antigens of, <u>Babesia</u>. The antigens used for this immunization can be those which are identified and described in the previous examples. The techniques employed to accomplish this immunization procedure are familiar to those skilled in this art. The spleens can then be removed from the immunized mice and the cells therefrom fused to SP-2 myeloma cells using polyethylene glycol. The desired hybrid cells can then be selected by adding hypozanthine-aminopterin-thymidine to the medium. The surviving cells can then be tested for antibody production. The testing for antibody production can be accomplished using IFA, ELISA, immunoblot, and/or immunoprecipitation procedures.

Example 21 - Detection of Babesia Antigens

The monoclonal antibodies, such as those produced by the procedure just described or those disclosed in Examples 1 and 9, can be used to test for the presence of <u>Babesia</u> antigens in a sample of biological fluid. Other monoclonal antibodies to <u>Babesia</u> antigens can also be used. The testing procedure involves contacting the biological fluid with a composition containing one or more of the monoclonal antibodies. If <u>Babesia</u> antigens are present in the biological fluid, then a reaction will occur and this reaction can be detected and quantified by fluorescence or other means.

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Example 22 - Detection of Anti-Babesia Antibodies

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Anti-<u>Babesia</u> antibodies can be detected in a fluid sample from a bovine suspected of containing these antibodies by performing ELISA procedures on the clinical samples. Generalized ELISA procedures are well known to those skilled in the art. The ELISA procedures or other simple diagnostic procedures of the subject invention could utilize as antigens, for example, whole cell or cell lysate using recombinant microorganisms which express <u>Babesia</u> antigens.

When the biological sample is contacted with the whole cell or cell lysate microorganisms, this contacting is done under conditions which will promote antigen/antibody immunocomplex formation between antigens expressed by the microorganism and antibodies present in the sample. The resulting immunocomplex can be readily detected utilizing standard labeling procedures.

It should be understood that the examples and embodiments described herein are for illustrative purposes only and that various modifications or changes in light thereof will be suggested to persons skilled in the art and are to be included within the spirit and purview of this application and the scope of the appended claims.



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BUDAPEST TREATY ON THE INTERNATIONAL RECOGNITION OF THE DEPOSIT OF MICROORGANISMS FOR THE PURPOSES OF PATENT PROCEDURE

INTERNATIONAL FORM

RECEIPT IN THE CASE OF AN ORIGINAL DEPOSIT ISSUED PURSUANT TO RULE 7.3 AND VIABILITY STATEMENT ISSUED PURSUANT TO RULE 10.2

To: (Name and Address of Depositor or Attorney)

Dr. Terry F. McElwain
Department of Veterinary Microbiology and Pathology
Washington State University
Pullman, Washington 99164-7040

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Deposited on Behalf of: Dr. Terry F. McElwain

Identification Reference by Depositor:

ATCC Designation

Hybridoma, 23.28.57.108

HB 10377

The deposit was accompanied by: ___ a scientific description ___ a proposed taxonomic description indicated above.

The deposit was received March 7, 1990 by this International Depository Authority and has been accepted.

AT YOUR REQUEST:

X We will not inform you of requests for the strain.

The strain will be made available if a patent office signatory to the Budapest Treaty certifies one's right to receive, or if a U.S. Patent is issued citing the strain.

If the culture should die or be destroyed during the effective term of the deposit, it shall be your responsibility to replace it with a living culture of the same.

The strain will be maintained for a period of at least 30 years after the date of deposit, and for a period of at least five years after the most recent request for a sample. The United States and many other countries are signatory to the Budapest Treaty.

The viability of the culture cited above was tested March 9, 1990. On that date, the culture was viable.

International Depository Authority: American Type Culture Collection, Rockville, Md. 20852 USA

Signature of person having authority to represent ATCC:

Bobbie A. Brandon, Head, ATCC Patent Depository

Date: March 12, 1990

cc: Roman Saliwanchik

Bruce Clary

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To: (Name and Address of Depositor or Attorney)

Terry F. McElwain, D.V.M., Ph.D. Washington State University Department of Veterinary Microbiology and Pathology Pullman, Washington 99164-7040

Deposited on Behalf of: Terry F. McElwain, D.V.M., Ph.D.

Identification Reference by Depositor:

ATCC Designation

Babesia bovis, Mexico Isolate

40601

The deposit was accompanied by: ___ a scientific description X a proposed taxonomic description indicated above.

The deposit was received May 3, 1989 by this International Depository Authority and has been accepted.

AT YOUR REQUEST:

X We will not inform you of requests for the strain.

The strain will be made available if a patent office signatory to the Budapest Treaty certifies one's right to receive, or if a U.S. Patent is issued citing the strain.

If the culture should die or be destroyed during the effective term of the deposit, it shall be your responsibility to replace it with a living culture of the same.

The strain will be maintained for a period of at least 30 years after the date of deposit, and for a period of at least five years after the most recent request for a sample. The United States and many other countries are signatory to the Budapest Treaty.

The viability of the culture cited above was tested <u>February 2, 1990</u>. On that date, the culture was viable.

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Signature of person having authority to represent ATCC:

Bobbie A. Brandon, Head, ATCC Patent Depository

Date: February 8, 1990

cc: Roman Saliwanchik Bruce Clary

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<u>Claims</u>

1	 A protein which has a molecular weight selected from the group
2	consisting of: 16, 25, 37, 42, 44, 55, 60, 85, 98, 125, 145, 225, and 250 kDa; and
3	which has one or more of the following characteristics:
4	(a) said protein elicits antibodies which bind to the surface of merozoites;
5	(b) said protein elecits an anti-Babesia immunologic response against
6	surface-exposed regions of merozoites; and
7	(c) said protein has merozoite-surface-exposed region(s) that can be
8	labeled with a surface-specific labeling procedure.
1	2. A Babesia merozoite surface protein having a molecular weight selected
2	from the group consisting of 16, 25, 37, 42, 44, 55, 60, 85, 98, 125, 145, 225, and 250
3	kDa.
1	3. The 42 kDa protein, according to claim 2, wherein said protein is a
2	glycosylated integral membrane protein.
1	4. A protein which has the amino acid sequence shown in Figure 1, or an
2	amino acid sequence which is substantially the same, or a fragment of the sequence
3	shown in Figure 1, so long as said substantially same sequence or said fragment
4	retains the biological activity of the sequence shown in Figure 1.
1	5. The protein, according to claim 4, which has the amino acid sequence
2	shown in Figure 1.
1	6. A Babesia bovis protein with species- and isolate-common epitopes
2	capable of eliciting immunologic responses in cattle, said protein having a molecular
3	weight selected from the group consisting of 19, 53, 59, and 120 kDa.

1	7. Monoclonal antibodies specific to a Babesia surface protein, where said
2	Babesia surface protein has a molecular weight selected from the group consisting
3	of: 16, 37, 42, 44, 60, 85, 145, and 225 kDa.
1	8. A monoclonal antibody selected from the group consisting of 23.8.34.24,
2	BABB75, MBOC79, 23.53.156, 23.38.120.8, 23.70.174.83, BABB35A ₄ , 23.3.16,
3	23.10.36, BABB93A ₁ , BABB90C ₄ , and 23.28.57.108.
1	9. A monoclonal antibody reagent useful in determining the presence of
2	pathogens, said reagent containing at least one monoclonal antibody species specific
3	to <u>Babesia</u> .
1	10. A Babesia DNA fragment that expresses a protein that reacts with an
2	antibody selected from the group consisting of: 23.8.34.24, BABB75, MBOC79,
3	23.53.156, 23.38.120.8, 23.70.174.83, BABB35A ₄ , 23.3.16, 23.10.36, BABB93A ₁ ,
4	BABB90C ₄ , and 23.28.57.108.
1	11. A <u>Babesia</u> DNA fragment which is selected from the group consisting
2	of lambda-Bo6, lambda-Bo25, lambda-Bo44-15, lambda-Bo44-16, lambda-Bo220-1,
3	and lambda-Bo220-2.
1	12. A DNA fragment comprising a 1.25 kb insert obtained by the digestion
2	of lambda-Bo44-15 or lambda-Bo44-16 with EcoRI.
1	13. The DNA sequence shown in Figure 1, or a DNA sequence which is
2	substantially the same, or a fragment of the sequence shown in Figure 1.
1	14. The DNA sequence shown in Figure 2, or a DNA sequence which is
2	substantially the same, or a fragment of the sequence shown in Figure 2.

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1	13. The DNA sequence snown in Figure 3, or a DNA sequence which is
2	substantially the same, or a fragment of the sequence shown in Figure 3.
1	16. A DNA sequence which codes for a polypeptide where said polypeptide
2	has the the amino acid sequence shown in Figure 1.
1	17. A DNA sequence which codes for a polypeptide having the amino acid
2	sequence shown in Figure 2.
1	18. A DNA sequence coding for the polypeptide shown in Figure 4.
1	19. A DNA sequence selected from the group consisting of
2	PQRPAETQQTQDSAAPSTPAAPSP and PQRPAETQQTQDSTAPGTPAAPSP,
3	and immunological equivalents thereof.
1	20. A recombinant microorganism transformed with lambda-Bo6, lambda-
2	Bo25, lambda-Bo220-1; lambda-Bo220-2; lambda-Bo44-15; lambda-Bo44-16; a 1.25
3	kb insert obtained by the digestion of lambda-Bo44-15 or lambda-Bo44-16 with
4	EcoRI; or a DNA sequence which is the same, substantially the same, or a fragment
5	of one of the sequences shown in Figure 1, Figure 2, or Figure 3.
1	21. A recombinant microorganism, according to claim 20, wherein said
2	microorganism is an Escherichia coli or Salmonella spp.
1	22. A recombinant protein produced by a microorganism transformed with
2	lambda-Bo44-15; lambda-Bo44-16; a 1.25 kb insert obtained by the digestion of
3	lambda-Bo44-15 or lambda-Bo44-16 with EcoRI; or a DNA sequence which is the
4	same, substantially the same, or a fragment of, one of the sequences shown in
5	Figure 1, Figure 2, or Figure 3.

1	23. The recombinant protein, according to claim 22, wherein said
2	recombinant protein has the amino acid sequence shown in Figure 1, Figure 2, or
3	Figure 4, or an amino acid sequence which is substantially the same, or a fragment
4	of one of those sequences.
1	24. A recombinant protein, according to claim 22, wherein said recombinant
2	protein elicits an antibody response to babesiosis in calves immunized with said
3	recombinant protein.
1	25. The use of a Babesia bovis protein having a molecular weight selected
2	from the group consisting of 16, 25, 37, 42, 44, 55, 60, 85, 98, 125, 145, 225, and 250
3	kDa, or an immunologic equivalent of said protein, to elicit neutralizing antibodies.
1	26. A vaccine for conferring immunity to babesiosis on a susceptible animal
2	host, said vaccine comprising an immunizing amount of bacterial cells, or products
3	of bacterial cells, where said cells have been transformed with DNA fragments from
4	Babesia that encode a surface protein.
1	27. A vaccine, according to claim 26, wherein said bacterial cells have been
2	transformed with lambda-Bo220-1; lambda-Bo220-2; lambda-Bo44-15; lambda-Bo44-
3	16; a 1.25 kb insert obtained by the digestion of lambda-Bo44-15 or lambda-B-44-16
1	with EcoRI; or a DNA sequence which is the same, substantially the same, or a
5	fragment of one of the sequences shown in Figure 1, Figure 2, or Figure 3.
1	28. A vaccine, according to claim 26, wherein said bacterial cells, or products
2	of bacterial cells, are in combination with an appropriate pharmaceutical carrier.
L	29. A vaccine comprising a protein, or mixture of proteins, where said
2	proteins are <u>Babesia</u> proteins, or immunologic equivalents thereof, where said

3	Babesia proteins have a molecular weight selected from the group consisting of 16,
4	25, 37, 42, 44, 55, 60, 85, 98, 125, 145, 225, and 250 kDa.
1	30. The vaccine, according to claim 29, wherein the protein, or at least one
2	of the proteins, has one of the amino acid sequences shown in Figure 1, Figure 2,
3	or Figure 4, or an amino acid sequence which is substantially the same, or a
4	fragment of one of those sequences.
1	31. A method for vaccinating a susceptible animal host to confer immunity
2	to babesiosis, said method comprising administering an immunizing amount of
3	bacterial cells, or products of bacterial cells, where said cells have been transformed
4	with DNA fragments from <u>Babesia</u> that encode a surface protein.
1	32. A method, according to claim 31, wherein said bacterial cells have been
2	transformed with lambda-Bo220-1; lambda-Bo220-2; lambda-Bo44-15; lambda-Bo44-
3	16; a 1.25 kb insert obtained by the digestion of lambda-Bo44-15 or lambda-B-44-16
4	with EcoRI; or a DNA sequence which is the same, substantially the same, or a
5	fragment of one of the sequences shown in Figure 1, Figure 2, or Figure 3.
1	33. A method, according to claim 31, wherein said bacterial cells, or
2	products of bacterial cells, are in combination with an appropriate pharmaceutical
3	carrier.
1	34. A method for detecting the presence of anti-Babesia antibodies in a
2	clinical sample of material suspected of containing these antibodies, said method
3	comprising the performance of a whole cell or cell lysate ELISA on the said clinical
4	sample, said ELISA is performed using recombinant microorganisms which express
5	Babesia specific antigens.

1	35. A method for detecting the presence of anti-Babesia antibodies in a
2	clinical sample suspected of containing these antibodies, said method comprising
3	(a) contacting the sample with whole cell or cell lysate recombinant
4	microorganisms which express Babesia specific antigens, said
5	contacting done under conditions which will promote specific
6	antigen/antibody immunocomplex formation between antigens
7	expressed by the microorganism and antibodies present in the sample;
8	and
9	(b) detecting immunocomplex formation by means of a label to thereby
10	detect the presence of <u>Babesia</u> antibodies in the sample.
1	36. A method of determining the presence of <u>Babesia</u> in a biological fluid,
2	said method comprising the steps of
3	(a) providing an aliquot of the biological fluid to be studied;
4	(b) contacting said aliquot with a measured amount of antibodies species
5	specific to Babesia antigens; and
6	(c) determining whether any reaction with said antibodies occurs.
1	37. A method, according to claim 36, wherein the antibodies have a
2	predetermined fluorescence response to a given optical stimulation.
1	38. A method, according to claim 36, wherein the amount of reaction is
2	determined and the presence of Babesia antigens is quantitatively derived
3	therefrom.
1	39. A method for the detection of evidence of babesiosis in bovine or tick
2	samples comprising tissue or fluid, said method comprising contacting said sample
3	with a DNA probe where said probe comprises labeled single-stranded DNA whose
4	sequence is sufficiently homologous with the DNA of Babesia so that the DNA of
5	the probe specifically binds with the DNA of said Babesia.

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1	40. A method for detecting evidence of Babesia bovis and/or Babesia
2	bigemina in a bovine or tick sample comprising tissue or fluid, said method
3	comprising contacting said sample with a DNA probe where said probe comprises
4	the DNA fragment known as lambda-Bo6, or an equivalent thereof.
1	41. A method of detecting evidence of Babesia bovis in a bovine or tick
2	sample comprising tissue or fluid, said method comprising contacting said sample
3	with a DNA probe where said probe comprises the DNA fragment lambda-Bo25,
4	or an equivalent thereof.
1	42. A plasmid selected from the group consisting of pBo6 and pBo25.
1	43. A method, according to claim 39, wherein said probe is labeled by means
2	of radioactive, fluorescent, bioluminescent, or chemiluminescent entities.
1	44. A method, according to claim 39, wherein said probe is labeled by biotin.
1	45. A probe for the detection of babesiosis, said probe comprising labeled
2	single-stranded DNA whose sequence is sufficiently homologous with the DNA of
3	Babesia so that the probe specifically binds with DNA of said Babesia.
L	46. A probe, according to claim 45, wherein said DNA comprises a Babesia
2	DNA which is the same, substantially the same, or a fragment of one of the DNA
3	sequences shown in Figure 1, Figure 2, or Figure 3.
l	47. A probe, according to claim 45, wherein said DNA comprises an insert
2	selected from the group consisting of lambda-Bo6, lambda-Bo25, lambda-Bo44-15,
}	lambda-Bo44-16, lambda-Bo220-1, and lambda-Bo220-2.

			10			20			3	0			40			50			60
SAA	TTC	AAT	BCT	TTT	CTT	8 AAT	GAC	AAT	£ TOO	000	CAT	ΔTG	\$ TTG	۵ΓΒ	ΔΔΤ	\$ CCC	۸۸۸	EAA	44
61u	Phe	Asn	Ala	Phe	Leu	Asn	Asp	Asn	Pro	Pro	His	Met	Leu	Thr	Asn	Gly	Lys	61u	Ly
			70			Bú			9				100						
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ATG Met	ACT The	GAA Blu	TAT	TAC	AAA	AAA	AAT	ATA	TCC	AAG	GAA	GAT	66T	BA6	GTA	AA6	GAT	TAC	AA
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			130 \$			140 t			15	0			160			170			180
ACT	ATG	GTC	AAG	TTT	TGC	AAC	GAT	m	CTA	GAC	AST	AAA	TCT	CCA	TTC	ATG	ASA	CTA	TAT
Thr	Het	Val	Lys	Phe	Cys	Asn	Asp	Phe	Leu	Asp	Ser	Ly5	Ser	Pro	Phe	Ħet	Arg	Leu	Tyr
			190			200			21	0			220			230			240
AAG	CAT	CTC	# AAT	844	TAT	‡ SAT	242	TΤΔ	‡ ata	200	AAC	۸۸۵	‡ *	ECA	PAA	\$	707	700	501
-75	His	767	Asn	Glu	Tyr	Asc	61u	Leu	Val	LVS	Lys	Lys	Pro	Aia	Gln	6lu	Ser	Ser	Pro
			250			260			270				280			290			
			1			\$			\$				ż			ż			300
6C: 4la	CCT Pro	TCA Ser	TCC Ser	CEG	CAA	AGA Aca	CCT	GCT Als	GAA	ACC	CAA	CAA	ACT	CA6	SAT	TCA	6CT	6CA	CCT
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AGC	4CT	232	GCA	G CT	CCT	TCA	CCC	cca	CAA	AGA	CCT	GCT	GAA	ACC	CAA	\$ Caa	ACT	CAG	GAT
Ser	Ihr	Pro	Ala	Ala	Pro	Ser	Pro	Pro	31n	Arg	Pro	Ala	6lu	Thr	Gln	Sln	Thr	Eln	Asp
		;	370			380			390)			100			410			420
TCA	ACT	RCA	CCT	RRC	ልቦተ	# 200	A79	GPT	‡	TCT	CCT	CAC	Ž CCA	rra.	ACT	\$		A 00	
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ICL Ser	CAA 61n	Ala	SAC Asp	CAC	CCA Pro	ACC Thr	AAA I ve	CCT	ACT The	CAG	ACA	CCT	GAA	66T	AAC	CTC	CAA	SGA	CAA
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CAG	GGT	ACA	ACC	AAB	CCA	939	GGA	TCT	TCA	TTC	400	TAT	660	SGA	TT6	ACT	8T6	939	ACT
61n	61 v	Thr	Thr	Lv5	Pro	Ala	Glv	Ger	Ser	Phe	Thr	Туг	Gly	Gly	Leu	Thr	Val	Ala	Thr
		;	550			560			570)		;	086			590			60Û
CTC	TGC	TAC	#	RTT	rtr	‡ 777	CLV	TTT	‡ TAG	TAA	PTA	ATC	‡	D+0		‡	~^~		*
Leu	Cvs	Tyr	TTC Phe	Val	Leu	Ser	Ala	?he			Leu	Het	Val	Val	Thr	Gin	186	Phe	Cvs
			510			620													
			1			1			6 <u>3</u> (‡				40 1			650 1			660 1
444	CTC	ATC	111	111	AAC	111	TAA	TGA	ATS	TTA	TGT	CTA	CAG	TAT	TCG	TSS	TAC	TTC	STC
LYS	Lan	ret	Phe	Phe	9sn	Phe			Yet	78.7	Cys	[51	Gln	Tyr	Ser	Trp	Tyr	Phe	Val

Figure 1 (continued)

6	70 8	980		690		700		710		720
AAT AGT ATA	GAC ATA	TCT ACA	ACA AGC	AGA GTA	CCA	4AA GTT	ATA	8 AGC AAT	ACT	8 STG CAC
Asn Ser Ile	Asp Ile	Ser Thr	Thr Ser	Arg Val	Pro	Lys Val	He	Ser Asn	Thr	Val His
7	/30 #	740 8		750 \$		760 1		770 \$		780
GGC ACA AAC	TGT TTG	TTA TAA	CTT TGT	COT TTO	ATT	TAA CAE	AAT	ACS TGG	TAT	STT ATA
6ly Thr Asn	tys Leu	ren	Leu Cys	Arg Phe	lle	Gln	Asn	Thr Trp	Tyr	Ile Leu
7	'90 ‡	800 t		810 \$		820 \$		830 1		840 \$
GAA TCT CAG	STA STT	CAT ATT	CAA TOT	CAA TEA	TA6	CTG ATT	AAT	TTG TTG	T66	ATA TAA
Glu Ser Gln		ure ite	ein CA2	61n		Leu Ile	Asn	Leu Leu	Trp	lle
	150 1	\$ 840		870 8		880 1		890 \$		900 1
CAG TGT GCG	ATT GCA	TTT CCA	TCT CTT	GTA ACA	776	ATA TTT	GAT	GGC ATC	AAA	CAG AGA
61n Cys Ala		rne Pro	ser Leu	vai inr	_61	ile Phe	Asp	Gly Ile	Lys	61n Arg
	10 \$	920 \$		930 8		940 \$		950 8		960 1
TAG CTG CAA	TAG GCT	AAG CAA	ACA AAA	GAA TGE	CTT	TAA TGG	AAA	TAA AGT	TAA	GCA TAT
Leu Gln	Ala	Lys Gin	Thr Lys	51u Trp	Leu	Trp	Lys	Ser		Ala Tyr
	70 \$	980 #		990 8		1000		1010 \$		1020
ATA TCG TAA	AAA TTA	AAA TAC	TAG CAT	ATT GCA	GAA	TAT AAA	CCT	6C6 TCT	GTT	TAA TTC
Ile Ser	LYS Leu	Lys iyr	His	lle Ala	Glu	Tyr Lys	Pro	Ala Ser	Val	Phe
10	30 #	1040 \$		1050 1		1060 1		1070 \$		1080 1
TTA ACS AAT	AAA AGT	GAA ATT	GTA ATA	CAC ATE	TAC	ATA CGA	TAG	ATG GGA	TAC	ATC CAC
Leu Thr Asn	Lys Ser	ein lie	Val Ile	His Net	Tyr	Ile Arg		Het Gly	Tyr	lle His
10	90 t	1100		1110 1		1120 1		1130 ‡		1140
ATG CGA CCA	TAT AAC	GAT STG	CGA ATT	TAG TAT	ATT	TAT ATC	AAA	CAT GAT	GAT	\$ 688 A66
Het Arg Pro	Tyr Asn	Asp Val	Arg Ile	Tyr	He	Tyr Ile	Lys	His Asp	Asp	Glu Arg
	50 1	1160		1170		1180		1190		1200
AAB GAT TAG	AAA ATC	AAC ATT	TAT AGT	# BCT CAA	TTA	TAT ATG	AAT	\$ GCA ATG	TTG	TEC TAC
Lvs Asp	Lys Ile	Asn Ile	Tyr Ser	Ala Sla	Leu	Tvr Met	Asn	Ala Met	Leu	Cys Tyr
	10	1220		1230						
ACA STT TST	‡ CAG AGG	1 CCT 6TC	AAT SIS	TAG AAT	TC				•	
Thr Val Cys	61n Arg	Pro Val	Asn Val	Asr	. •					
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Figure 2

PBV42 NUCLEOTIDE SEQUENCE

1	•	СТТСА	ስ ጥሮርጥሮርጥጥር	` <i>CCC</i> III <i>CC</i> III	ATTCTACGAT
36	GACATGTCTA				
86					AGACCAAATT
					GATGATCAGA
136	AGGTAAAAGA	CACATTCAAA	AATTTATACA	AAGTCAACGC	ATTGATAAAG
186	AACAATCCTA	TGATTCGCCC	TGATCTATTT	' AATGCAACTA	TTGTTAGCGG
236	TTTTTCAACT	AAGAATGACG	AGGAAAAATT	CAATGCTATA	TTTGATTCCA
286	TTAAGGGAAT	GTACTATAGA	GCTCAACACA	TGGACAAATA	TTTGAAGTCA
336					AGGCAGTTGA
386	ATATTTCAAG	AAGCATGTTT	ATACGGGGGA	ACACGTTGTT	GACGTCAACG
436	GTATGGCTGG	TGTTTGCAAG	GAGTTTTTAA	GCCCGGCCTC	TGATTTCTAC
486	AAACTTGTTG	AGTCTTTTGA	TGCGTTTGCA	CATGCTAAGG	TGCACGCTCA
536	AGTAGGAAAT	TTTGTTAAAC	CTGGAACTGA	CATCGCTCCT	CCTAAGGATG
586	TTACTGATGC	ATTAGAAAAG	GAATTGCAAG	AGCAAAAACC	TGCACGAAGT
636	GAGAGCACCG	AAGTACCCGC	TCCAGGTGAT	GCATCTGGCG	TCCAACAACC
686	GCCTGCATCA	GGAACATCCC	CGCAAGGACC	TGCTCCGACT	ACACCCAGCC
736	CATCTCCAGA	GTCCTCAGGA	AACCTCCAAG	GACAACAGGG	TACAACCAAG
786	CCAGCCGGAT	CTTCTTTCAC	CTATGGCGGA	TTGACTGTGG	CTACTCTCTG
836	CTACTTCGTT	CTCTCTGCAT	TTTAAAAACT	AATGGTAGTG	ACACAATAGT
886	TTTGTAAACT	CATGTTTTTT	AACTTTTAAT	ATGTAGTGAA	AAA
TRANSLATI	ON ofrBv42				
1	SIVLP	EGSFYDDMSK	FYGAVGSFDQ	TKLYSVLSAN	FKAAKMDDQK
46	VKDTFKNLYK	VNALIKNNPM	IRPDLFNATI	VSGFSTKNDE	EKFNAIFDSI
96	KGMYYRAQHM	DKYLKSLRWN	TDIVEEDREK	AVEYFKKHVY	TGEHVVDVNG
146	MAGVCKEFLS				
196					
246					

Figure 3

1 GACGGATAGT ATTTTACATA TACATTTGTC GACTTTTATA TATAGCAGTG CTATAGACAA ACAATACACA GATTAATCTT TAGATACTAA STICAATAAT 101 ATTACGGACA TATTGTAGAC AATGAGAATC ATTAGCGGCG TTGTCGGTTG CCTTTTCTTG GTGTTTTCAC ACCATCTGTC TGCTTTTCGC CACAATCAGA 151 BAGTAGGAAG TETEGETECA GETGANITGG TAGGTGATTT AACETECACA TTGGAAACAG CTGATACTTT GATGACTCTC DGTGACCACA TGCACACAT 301 TACTAAGGAT ATGAAACATG TTTTGAGCAA TGGTCGTGAB CAGATTGTAA ATGATGTTTG CTCTAATGCT CCTGAGGACT CCAACTGTCS TGAUGTAGTT 331 RACAATTATE CTEACCETTE TEAHATETAC ESAIGCITTA CSATTSACAA TETCHARTAT COSTTETATO ARGENTACCA ACCIOTATOT CITTECHARCO CTTACCASTT GGATCHTGCS TTCAGATTGT TCGAGGAGAG TTCATCGAGC 501 CCTGCCAAGA ACAGCGTAAA ACGGGAAFICS ITGCGTTTC: GAAATGGAGC GAACCATEGI GATTACCACT ACTTCCICAC TEGTCTGTTG PACAACAATG TIGIBCACGA GGAAGGAACT ACCGAIGITS AATAICTIGI CAACAAGGTA 551 CTCTATATGG CTACCATGAG CTACAGAGT TATTTGACAG TAACAGTAT GANCGCCAAG TTCTTCAACA GATTCAGCTT CACTACAAAG ATGTTCAGTC GTCGTATTAG GCAAACATTG AGTGATATCA TCAGGTGGAA TSTTCCTGAA 801 GATTTTGAAG AAARGAGCAT CGAACGTATC ACTCAACTTA CTAGCAGCTA CSAAGATTAC AIGTTENECE AGATTCCPAC TCTTTCCAAG TTTGCACGTC 751 GTTATGCTGA CATGGTGAAG AAGGTTCTGC TCGGTAGCTT GACCTCGTAC STTGAAGCTC CTTGETACAA AAGATGGATA AAGAAATTCA GAGACTTTTT 1951 CTETABBAAC GTTACCCAAC CTACAAGAA GTTCATCGAG GATACTAACG ANGITACCAA ANACTATETS AAAGCCAATE TIGCTGNGCC CACTAAAAAG 1101 TTTATGCAGG ACACTCACGA AAAAACCAGA GGCTATCTGA AAGAGAATGT 1151 1201 AGCCGRACCT ACTAAGACTT TTTTCAAGGA GGCTCCTCAA GTCACCAAAC 1251 ACTITITUGA TGAGAACATT GGCCAACCCA CCAAGGAGTT TTTCAGGGAA 1301 SCTCCCCAAG CCACTAGACA TTTCCTAGAC GAAAACATCG GTCAACCAAC 1351 CAAGGAGTTC TTCAGGGAGG CTCCTCAAGC CACTAAGCAC TTCCTAGGCG AGASTATTEC TEAACCTACT AAAGAATITI TCAAGGATGT CCCTCAAGTC 1401 ACCAAGAAGG TTATAACTGA GAACATTGCT CAACCAACTA AGGAGTTCCG GAGGGAGGTT CCTCATGCTA CCATGAAAGT CTTGAATGAA AACATTGCTC 1501 AACCTGCCAA GGAAATCATA CATGAGTTTG GTACAGGCGC CAAGAATTTC 1551 1601 ATTTECECAG CECATGAAGG TACTAAGCAG TTETTAAACG AAACTGTTGG CCAACCTACA AAGGAATTCC TGAACGGAGC TTTAGAAGACT ACTAGAGACG CATTACACCA TCTGGGTAAA TCATCAGAAB AAGCCAACCT TTATGATGCC 1751 ACGGAAAATA CCACTCAGGC TAACGACTCA ACTACTTCCA ACGGTGAAGA CACCECEGGA TACCTCTGAT GAGATECETT TATAATEGCA CAAACTCAAC 1851 AAATGATGTA TCGTCATCTG ATCCATCGGT TTTCAATATT GTATTGGATG 1901 CAATATCTGA ATGCATATGA TGCGACAGTT TCCATCATCG GGTGCCGAAT 1951 CGTAACTCTC ATAACACCAT TITAAGTTAT SCTCGTGCCG

Figure 4

MRIISGYVGC LFLVFSHHVS AFRHNORVGS LAPAEYVGDL TSTLETADTL MTLRDHMHNI TKDMKHVLSN GREQIVNDVC SNAPEDSNCR EVVNNYADRC 51 EMYGCFTIDN VKYPLYGEYG PLSLPNPYGL DAAFRLFKES ASNPAKNSVK 101 151 REWLRFRNGA NHGDYHYFVT GLLNNNVVHE EGTTDVEYLV NKVLYMATMN 201 YKTYLTVNSM NAKFFNRFSF TTKIFSRRIR QTLSDIIRWN VPEDFEERSI ERITQLTSSY EDYMLTQIPT LSKFARRYAD MVKKVLLGSL TSYVEAPWYK 251 RWIKKFROFF SKNVTQPTKK FIEDTNEVTK NYLKANVAEP TKKFMQDTHE 301 KTKGYLKENV AEPTKTFFKE APQVTKHFFD ÉNIGQPTKEF FREAPQATKH 351 FLDENIGOPT KEFFREAPQA TKHFLGENIA QPTKEFFKDV PQVTKKVITE 401 NIAQPTKEFR REVPHATMKV LNENIAQPAK EIIHEFGTGA KNFISAAHEG 451 TKOFLNETVG OPTKEFLNGA LETTKDALHH LGKSSESANL YDATENTTQA 501 551 NDSTTSNGED TAGYL

INTERNATIONAL SEARCH REPORT

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FURTHER INFORMATION CONTINUED FROM THE SECOND SHEET	
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Claims 25, 31 - 33	
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